OPE

CRF Problem Report

The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: _09/621, 630_

Filing Date: _7/21/2000_

Date Processed by STIC: _7/31/2000_

STIC Contact: **Mark Spencer, 703-308-4212**

Nature of Problem:

The CRF (was):

_____ (circle one) Damaged or Unreadable (for Unreadable, see attached)

_____ Blank (no files on CRF) (see attached)

_____ Empty file (filename present, but no bytes in file) (see attached)

_____ Virus-infected. Virus name: _____ The STIC will not process the CRF.

_____ Not saved in ASCII text

_____ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.

__✓__ Did not contain a Sequence Listing. (see attached sample)

_____ Other:

_____

PLEASE USE THE CHECKER VERSION 3.0 PROGRAM TO REDUCE ERRORS. SEE BELOW FOR DETAILS:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
http://www.uspto.gov/web/offices/pac/checker

SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES ENCODED THEREBY

This application claims priority under 35 USC §119(e), §119(a-d) and §120 of the following applications, the entire contents of which are hereby incorporated by reference:

Country	Filing Date	Attorney No.	Client No.	Application No.
United States	07/21/99	2750-0503P	00066.001	60/144,814

5

FIELD OF THE INVENTION

The present invention relates to isolated polynucleotides that represent a complete

sample of submitted CD file

Per 1.824 of Sequence Rules, submit only a complete Sequence Listing

Amended Compact Discs

THE ATTACHED CD (COPY 1) HAS BEEN REVIEWED BY OIPE FOR COMPLIANCE WITH 37 CFR 1.52(E).

Date: _4/15/02_

Serial No./Control No. _09 621 630_

Reviewed By: _Williams_   Phone: _305 3021_

☑ The compact discs are readable and acceptable.

☐ Copy 1 and Copy 2 of the compact discs are not the same.

☐ The compact discs are unreadable.

☐ The files on the compact discs are not in ASCII.

☐ The compact discs contain at least one virus.

☐ Other

_____

_____

_____

<110> ALEXANDROV, Nickolai
      TROUKHAN, Maxim

<120>   Sequence-Determined DNA Fragments and Corresponding Polypeptides Encoded Thereby

<130>   2750-0991P

<140>   09/621,630
<141>   2000-07-21

<160>   769

<170>   PatentIn version 3.0

<210>   1
<211>   5
<212>   PRT
<213>   Artificial Sequence

<220>
<223>   Consensus Pattern from various organisms

<400>   1

Leu Ile Val Met Thr
1   5

<210>   2
<211>   5
<212>   PRT
<213>   Artificial Sequence

<220>
<223>   Consensus Pattern from various organisms

<400>   2

Leu Ile Val Met Phe
1   5

<210>   3
<211>   5
<212>   PRT
<213>   Artificial Sequence

<220>
<223>   Consensus Pattern from various organisms

<400>   3

Gly Ala Thr Met Cys
1   5

<210>   4
<211>   4
<212>   PRT
<213>   Artificial Sequence

<220>
<223>   Consensus Pattern from various organisms

```
Volume in drive D:\ is 2750-0991P
Directory of D:\

new2750-0991P.ST25.txt                          148 KB      4/3/02


  1 file(s)
  Total filesize 148 KB
  0 kilobytes free
```